IN THE CLAIMS:

Please cancel claims 2, 10, 12 and 24-28 without prejudice or disclaimer.

1. (Thrice Amended) A method for producing a <u>recombinant</u> filamentous <u>Actinomycete</u> bacterium exhibiting reduced branching and fragment septation during growth, said method comprising:

providing a filamentous <u>Actinomycete</u> bacterium, said filamentous <u>Actinomycete</u> bacterium lacking <u>significant detectable</u> endogenous <u>ssgA-activity SsgA</u>, with the capability of having or expressing heterologous <u>SsgA-activity SsgA</u>, which <u>activity heterologous SsgA</u>, in <u>Streptomyces griseus</u>, is encoded by an ssgA gene <u>having at least comprising</u> the sequence:



2. Canceled.

- 3. (Twice amended) The method according to claim 1, wherein said additional heterologous SsgA-activity SsgA is provided by transfecting or transforming said filamentous Actinomycete bacterium with additional genetic information DNA encoding said activity SsgA.
- 4. (Amended) The method according to claim 3, wherein said additional genetic information <u>DNA</u> comprises an <u>the</u> ssgA gene or a derivative or fragment thereof encoding similar SsgA-activity.

- 5. (Amended) The method according to claim 4, wherein said ssgA gene is derived from of an actinomycete origin.
- 6. (Amended) The method according to claim 4, wherein said <u>ssgA</u> gene is <u>derived</u> from of a streptomycete origin.
- 7. (Amended) The method according to claim 5, wherein said <u>ssgA</u> gene is <u>derived</u> from <u>of Streptomyces Streptomyces</u> griseus, Streptomyces collinus, Streptomyces albus, Streptomyces goldeniensis or Streptomyces netropsis <u>origin</u>.
- 8. (Twice amended) The method according to claim 3, wherein said additional genetic information <u>DNA</u> is integrated into the bacterial genome of the filamentous Actinomycete bacterium.
- 9. (Twice amended) The method according to claim 3, wherein said additional genetic information DNA is part of an episomal element.
 - 10. Canceled.
- 11. (Twice amended) The method according to claim 3 wherein <u>expression of said ssgA-activity SsgA</u> is inducible or repressible with a signal.
 - 12. Canceled.
- 13. (Amended) The method according to claim 12 3, wherein said <u>filamentous</u> Actinomycete bacterium is a Streptomyces.
- 14. (Twice amended) The method according to claim 3 wherein said filamentous Actinomycete bacterium produces a useful product.

- 15. (Original) The method according to claim 14 wherein said useful product is an antibiotic.
- 16. (Original) The method according to claim 14, wherein said useful product is a protein.
- 17. (Amended) The method according to claim 16 wherein said protein is heterologous to said filamentous Actinomycete bacterium.
- 18. (Twice amended) The method according to claim 16, wherein said protein is expressed from a vector encoding said protein present in said filamentous <u>Actinomycete</u> bacterium.
- 19. (Twice amended) The method according to claim 18, wherein said protein is secreted by said filamentous Actinomycete bacterium.

24-28. Canceled.

- 29. (New) The method according to claim 1, wherein the ssgA gene encodes a protein comprising SEQ ID NO: 3.
- 30. (New) A method for producing a recombinant Actinomycete bacterium, said method comprising:

transforming an Actinomycete bacterium lacking a detectable endogenous SsgA with a means for enhancing septation and fragmentation in a culture of the Actinomycete bacterium;

wherein the Actinomycete bacterium is selected from the group consisting of Streptomyces coelicolor, Streptomyces lividans, Streptomyces clavuligerus and Saccharopolyspora erythraea.

- 31. (New) The method according to claim 30, wherein the means for enhancing septation and fragmentation comprises SEQ ID NO: 1.
- 32. (New) The method according to claim 30, wherein the means for enhancing septation and fragmentation encodes a protein comprising SEQ ID NO: 3.
- 33. (New) A method for producing a recombinant Actinomycete bacterium, said method comprising:

transforming an Actinomycete bacterium lacking a detectable endogenous SsgA with a nucleic acid encoding a heterologous SsgA comprising SEQ ID NO: 3.

34. (New) The method according to claim 33, wherein Actinomycete bacterium is selected from the group consisting of *Streptomyces coelicolor*, *Streptomyces lividans*, *Streptomyces clavuligerus* and *Saccharopolyspora erythraea*.